

INFORMATION DISCLOSURE

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APPLICANT
Thomas Charles ELLEMAN, et al.

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**GROUP
1631**

U.S. PATENT DOCUMENTS

[illegible]

FOREIGN PATENT DOCUMENTS

EXAMINER'S INITIALS	CITE NO.	Foreign Patent Document Country Codes - Number 4 - Kind Codes (if known)	Publication Date MM-DD-YYYY	Name of Patentee or Applicant of Cited Document	Pages, Columns, Lines Where Relevant Figures Appear	Translation	
						Yes	No
		WO 99/28347	06/10/1999	COMMONWEALTH SCIENTIFIC AND INDUSTRIAL RESEARCH ORGANISATION		X	

OTHER ART (Including Author, Title, Date, Pertinent Pages, Etc.)

EXAMINER'S INITIALS	CITE NO.	Include name of the author (in CAPITAL LETTERS), title of the article (when appropriate), title of the item (book, magazine, journal, serial, symposium, catalog, etc.), date, page(s), volume-issue number(s), publisher, city and/or country where published.	
MPA		SINGH, Juswinder., et al. "Structure-Based Design of a Patent, Selective and Irreversible Inhibitor of the Catalytic Domain of the erbB Receptor Subfamily of Protein Tyrosine Kinases." Journal of Medicinal Chemistry, American Chemical Society, Volume 40, March 28, 1997, pp. 1130-1135, XP002107337	
↓		GARRETT, Thomas P., et al. "Crystal structure of the first three domains of the type-1 Insulin-like growth factor receptor." Nature (London) Volume 394, Number 6691, July 23, 1998, pp.395-399, XP002297751	
		MCINNIS, Cambell., et al. "NMR study of the transforming growth factor-alpha (TGF-alpha)-epidermal growth factor receptor complex. Visualization of human TGF-alpha binding determinants through nuclear Overhauser enhancement analysis." Journal of Biological Chemistry, Volume 271, Number 50, 1996, pp.32204-32211, XP002297752	
		SANCHEZ, Roberto., et al. "Evaluation of Comparative Protein Structure Modeling by MODELLER-3" Proteins, Structure, Function and Genetics, Suppl. Pp. 1:50-58 (1997	

EXAMINER

DATE CONSIDERED

EXAMINER
Marianne P. Allen

12/23/04

*EXAMINER: Initial if reference considered, whether or not citation is in conformance with MPEP 609. Draw line through citation if not in conformance and not considered.

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